

## RAW SEQUENCE LISTING

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The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/511,327  
Source: PUT/10  
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## RAW SEQUENCE LISTING

DATE: 05/05/2005

PATENT APPLICATION: US/10/511,327

TIME: 10:29:27

Input Set : A:\2005-01-19 0933-0230PUS1.ST25.txt

Output Set: N:\CRF4\05052005\J511327.raw

3 <110> APPLICANT: Harri SAVILAHTI et al.  
 5 <120> TITLE OF INVENTION: METHOD AND MATERIALS FOR PRODUCING DELETION DERIVATIVES OF  
 POLYPEPTIDES

7 <130> FILE REFERENCE: 0933-0230PUS1  
 9 <140> CURRENT APPLICATION NUMBER: US 10/511,327  
 10 <141> CURRENT FILING DATE: 2004-10-15  
 12 <160> NUMBER OF SEQ ID NOS: 16  
 14 <170> SOFTWARE: PatentIn Ver. 2.1  
 16 <210> SEQ ID NO: 1  
 17 <211> LENGTH: 54  
 18 <212> TYPE: DNA  
 19 <213> ORGANISM: Artificial Sequence  
 21 <220> FEATURE:  
 22 <223> OTHER INFORMATION: Description of Artificial Sequence: Modified Mu  
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 32 <213> ORGANISM: Artificial Sequence  
 34 <220> FEATURE:  
 35 <223> OTHER INFORMATION: Description of Artificial Sequence: Modified Mu  
 36 transposon  
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 41 agcacacggt cacactgctt ccggtagtag ataaaccggt aaaccagcaa tagacataag 180  
 42 cggctattta acgaccctgc cctgaaccga cgaccgggtc gaatttgctt tcgaatttct 240  
 43 gccattcatc cgcttattat cacttattca ggcgtagcaa ccaggcggtt aagggcacca 300  
 44 ataactgcct taaaaaaatt acgccccgcc ctgccactca tcgcagtact gttgtaattc 360  
 45 attagcatt ctgccgacat ggaagccatc aaaaacggca tgatgaacct gaatcgccag 420  
 46 cggcatcagc accttgctgc cttgcgtata atatttgccc atggtgaaaa cgggggagcga 480  
 47 gaagtgtcc atattggcca cgtttaaatc aaaactggtg aaactcacc agggattggc 540  
 48 tgagacgaaa aacatattct caataaacc tttagggaaa taggccagg tttcaccgta 600  
 49 acacgccaca tcttgccaat atatgtgtag aaactgccgg aaatcgctcg ggtattcact 660  
 50 ccagagcgat gaaaacggtt cagtttgctc atggaaaacg gtgtaacaag ggtgaacact 720  
 51 atcccatatc accagctcac cgtctttcat tgccatacgt aattccggat gagcattcat 780  
 52 caggcgggca agaattgtga taaaggccgg ataaaacttg tgcttatttt tctttacggt 840  
 53 ctttaaaaag gccgtaatat ccagctgaac ggtctgggta taggtacatt gagcaactga 900  
 54 ctgaaatgcc tcaaaatggt ctttacgatg ccattgggat atatcaacgg tggatatatcc 960  
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 57 aacgtctcat ttccgcaaaa agttggccca gggcttcccg gtatcaacag ggacaccagg 1140

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58 atttatttat tctgcgaagt gatcttccgt cacaggtatt tattcggtcg aaaaggatcc 1200
59 gttttcgcatt ttatcgtgaa acgctttcgc gtttttcggt caatcaatca gatc      1254
62 <210> SEQ ID NO: 3
63 <211> LENGTH: 54
64 <212> TYPE: DNA
65 <213> ORGANISM: Bacteriophage Mu
67 <400> SEQUENCE: 3
68 gatctgaagc ggcgcacgaa aaacgcgaaa gcgtttcacg ataaatgcga aaac      54
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76 <220> FEATURE:
77 <223> OTHER INFORMATION: Description of Artificial Sequence: Modified Mu
78     end sequence
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85 <211> LENGTH: 50
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87 <213> ORGANISM: Artificial Sequence
89 <220> FEATURE:
90 <223> OTHER INFORMATION: Description of Artificial Sequence: Modified Mu
91     end sequence without 5' overhang
93 <400> SEQUENCE: 5
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100 <213> ORGANISM: Artificial Sequence
102 <220> FEATURE:
103 <223> OTHER INFORMATION: Description of Artificial Sequence: Sequencing
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106 <400> SEQUENCE: 6
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111 <211> LENGTH: 4814
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115 <220> FEATURE:
116 <223> OTHER INFORMATION: Description of Artificial Sequence: Modified Tn7
117     transposon
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121 ccagcagccg cagcggcgcg atctcgggca gcgttgggtc ctggccacgg gtgcgcatga 120
122 tcgtgctcct gtcgttgagg acccggttag gctggcgggg ttgccttact ggtagcaga 180
123 atgaatcacc gatacgcgag cgaacgtgaa gcgactgctg ctgcaaaacg tctgcgacct 240
124 gagcaacaac atgaatggtc ttcggtttcc gtgtttcgta aagtctggaa acgcggaagt 300
125 cagcgccctg caccattatg ttccggatct atgtcgggtg cggagaaaga ggtaatgaaa 360

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127 atattctttt ttttcttata aaacttaaaa ccttagaggc tatttaagtt gctgatttat 480
128 attaatTTTA ttgttcaaac atgagagctt agtacgtgaa acatgagagc ttagtacggt 540
129 agccatgaga gcttagtacg ttagccatga gggtttagtt cgTTaaacat gagagcttag 600
130 tacgtTaaac atgagagctt agtacgtgaa acatgagagc ttagtacgta ctatcaacag 660
131 gttgaactgc tgatcttcgg atctatgtcg ggtgcggaga aagaggtaat gaaatggcag 720
132 atccctggct tgttgTccac aaccgtTaaa cctTaaaagc tTaaaagcc ttatatattc 780
133 ttttttttct tataaaactt aaaaccttag aggctattta agttgctgat ttatattaat 840
134 tttattgttc aaacatgaga gcttagtacg tgaacatga gagcttagta cgTtagccat 900
135 gagagcttag tacgttagcc atgagggttt agttcgTtaa acatgagagc ttagtacggt 960
136 aaacatgaga gcttagtacg tgaacatga gagcttagta cgtactatca acaggTgaa 1020
137 ctgctgatct tcggatctat gtcgggtgcg gagaaaggg taatgaaatg gcatccggat 1080
138 ctgcatcgca ggatgtgct ggtaccctg tggaaacacct acatctgtat taacgaagca 1140
139 ttattgaagc atttatcagg gttattgtct catgagcgga tacatatTTg aatgtattta 1200
140 gaaaaataaa caaatagggg ttccgcgcac atttccccga aaagtgccac ctgacgtcta 1260
141 agaaaccatt attatcatga cattaacctt taaaaatagg cgtatcacga ggccctttcg 1320
142 tcttcaagaa ttctcatggt tgacagctta tcatcgataa gctTtaatgc ggtagtttat 1380
143 cacagtTaaa ttgctaacgc agtcaggcac cgtgtatgaa atctaacaat gcgctcatcg 1440
144 tcatcctcg caccgtcacc ctggatgtcg taggcattag cttggTtatg ccggtactgc 1500
145 cgggcctctt gcgggatatc gtccattccg acagcatcg cagtcactat ggcgtgctgc 1560
146 tagcgctata tgcgtgatg caatttctat gcgcaccctg tctcgagca ctgtccgacc 1620
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148 tcatggcgac cacaccctgc ctgtggatcc tctacgccg acgcatcgTg gccggcatca 1740
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152 tcaacggcct caacctacta ctgggctgct tctaatgca ggagtcgat aaggagagc 1980
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158 tgggctacgt cttgctggcg ttccgcgacgc gaggtggat ggcttcccc attatgattc 2340
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175 ggccgctaac tataacggtc ctaaggtagc gagtttaaac gatatcggat ccggccgccc 3360
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177 atccagccag aaagtgaggg agccacggtt gatgagagct ttgtttaggagg tggaccagtt 3480
178 ggtgattttg aacttttgct ttgccacgga acggtctgcg ttgtcgggaa gatgcgtgat 3540
179 ctgatccttc aactcagcaa gagttcgatt tattcaacaa agccgcccgtc ccgtcaagtc 3600
180 agcgtaatgc tctgccagtg ttacaaccaa ttaaccaatt ctgattagaa aaactcatcg 3660
181 agcatcaaat gaaactgcaa ttatttcata tcaggattat caataccata tttttgaaaa 3720
182 agccgtttct gtaatgaagg agaaaactca ccgaggcagt tccataggat ggcaagatcc 3780
183 tggatcgggt ctgcgattcc gactcgtcca acatcaatac aacctattaa tttcccctcg 3840
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187 aatacgcgat cgctgtttaa aggacaatta caaacaggaa tcgaatgcaa ccggcgagag 4080
188 aacactgccg gcgcatcaac aatattttca cctgaatcag gatattcttc taatacctgg 4140
189 aatgctgttt tcccggggat cgagtggtg agtaaccatg catcatcagg agtacggata 4200
190 aaatgcttga tggtcggaag aggcataaat tccgtcagcc agtttagtct gaccatctca 4260
191 tctgtaacat cattggcaac gctacctttg ccatgtttca gaaacaactc tggcgcatcg 4320
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194 tcccggtgaa tatggctcat aacaccctt gtattactgt ttatgtaagc agacagtttt 4500
195 attgttcatg atgatattt tttatcttgt gcaatgtaac atcagagatt ttgagacaca 4560
196 acgtggctta ctaggatccg atatcattta aatctaggga taacagggta atactagtgt 4620
197 cgaccaacca gataagtga atctagttcc aaactatttt gtcattttta attttcgtat 4680
198 tagcttacga cgctacaccc agttcccatc tattttgtca ctcttcccta aataatcctt 4740
199 aaaaactcca tttccacccc tcccagttcc caactatttt ctactcactc accgtaagat 4800
200 gcttttctgt gact 4814
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212 <400> SEQUENCE: 8
213 acggtgagtg agtagaaaat agttgggaac tggga 35
216 <210> SEQ ID NO: 9
217 <211> LENGTH: 43
218 <212> TYPE: DNA
219 <213> ORGANISM: Artificial Sequence
221 <220> FEATURE:
222 <223> OTHER INFORMATION: Description of Artificial Sequence:
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225 <400> SEQUENCE: 9
226 cgtatgagtg agtagaataa agtcttaaac tgaacaaaat aga 43
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232 <213> ORGANISM: Artificial Sequence
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256 <211> LENGTH: 21
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258 <213> ORGANISM: Artificial Sequence
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262     Oligonucleotide primer
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265 agctggcgaa agggggatgt g                                21
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271 <213> ORGANISM: Artificial Sequence
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274 <223> OTHER INFORMATION: Description of Artificial Sequence:
275     Oligonucleotide primer
277 <400> SEQUENCE: 13
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281 <210> SEQ ID NO: 14
282 <211> LENGTH: 50
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284 <213> ORGANISM: Bacteriophage Mu
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291 <211> LENGTH: 50
292 <212> TYPE: DNA
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296 <223> OTHER INFORMATION: Description of Artificial Sequence: Modified Mu end sequence
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303 <211> LENGTH: 50
304 <212> TYPE: DNA
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